

1653

RAW SEQUENCE LISTING
PATENT APPLICATION: US/08/962,560B
DATE: 09/14/2000
TIME: 17:33:46
Input Set : A:\PTO.txt
Output Set: N:\CRF3\09142000\H962560B.raw

Does Not Comply
Corrected Diskette Needed

See pp. 1, 2, 3

4 <110> APPLICANT: Hilton, Douglas J.
5 Alexander, Warren S.
6 Viney, Elizabeth M.
7 Wilson, Tracy A.
8 Richardson, Rachel
9 Starr, Robyn
10 Nicholson, Sandra E.
11 Metcalf, Donald
12 Nicola, Nicos A.
14 <120> TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC AGENTS
16 <130> FILE REFERENCE: Davies Collision Cave
18 <140> CURRENT APPLICATION NUMBER: 08/962,560B
19 <141> CURRENT FILING DATE: 1997-10-31
21 <160> NUMBER OF SEQ ID NOS: 68
23 <170> SOFTWARE: Patentin Ver. 2.1

ERRORED SEQUENCES

137 <210> SEQ ID NO: 4
138 <211> LENGTH: 212
139 <212> TYPE: PRT
140 <213> ORGANISM: Mus musculus
142 <400> SEQUENCE: 4
143 Met Val Ala Arg Asn Gln Val Ala Ala Asp Asn Ala Ile Ser Pro Ala
144 1 5 10 15
146 Ala Glu Pro Arg Arg Ser Glu Pro Ser Ser Ser Ser Ser Ser
147 20 25 30
149 Ser Pro Ala Ala Pro Val Arg Pro Arg Pro Cys Pro Ala Val Pro Ala
150 35 40 45
152 Pro Ala Pro Gly Asp Thr His Phe Arg Thr Phe Arg Ser His Ser Asp
153 50 55 60
155 Tyr Arg Arg Ile Thr Arg Thr Ser Ala Leu Leu Asp Ala Cys Gly Phe
156 65 70 75 80
158 Tyr Trp Gly Pro Leu Ser Val His Gly Ala His Glu Arg Leu Arg Ala
159 85 90 95
E--> 161 Glu Pro Val Gly Thr Phe Leu Val Arg Asp Ser Arg Gln Arg Asn Cys
162 100 105 110
164 Phe Phe Ala Leu Ser Val Lys Met Ala Ser Gly Pro Thr Ser Ile Arg
165 115 120 125
167 Val His Phe Gln Ala Gly Arg Phe His Leu Asp Gly Ser Arg Glu Thr
168 130 135 140
171 Phe Asp Cys Leu Phe Glu Leu Leu Glu His Tyr Val Ala Ala Pro Arg
172 145 150 155 160
174 Arg Met Leu Gly Ala Pro Leu Arg Gln Arg Arg Val Arg Pro Leu Gln
175 165 170 175
177 Glu Leu Cys Arg Gln Arg Ile Val Ala Ala Val Gly Arg Glu Asn Leu

Invalid amino acid
designator.

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178          180          185          190
180 Ala Arg Ile Pro Leu Asn Pro Val Leu Arg Asp Tyr Leu Ser Ser Phe
181          195          200          205
183 Pro Phe Gln Ile
184          210
1644 <210> SEQ ID NO: 24
1645 <211> LENGTH: 2019
1646 <212> TYPE: DNA
1647 <213> ORGANISM: Mus musculus
1649 <220> FEATURE:
1650 <221> NAME/KEY: UNSURE
1651 <222> LOCATION: (1981)
1652 <223> OTHER INFORMATION: N is unsure
1654 <220> FEATURE:
1655 <221> NAME/KEY: UNSURE
1656 <222> LOCATION: (1992)
1657 <223> OTHER INFORMATION: N is unsure
1659 <220> FEATURE:
1660 <221> NAME/KEY: UNSURE
1661 <222> LOCATION: (2010) → n is at location 2000
1662 <223> OTHER INFORMATION: N is unsure
1664 <400> SEQUENCE: 24
1665 ggcacgaggc ggggtcaggg cggaggtcga ggaccaagta ggcattggcg agggcgggac 60
1666 cggcccgcat ggacggggcg gcccgggacc cgcaggtcct aatctgaagg agtggctgag 120
1667 ggagcagttc tctgaccatc cactggagca ctgtgacgat acaagactcc atgatgcagc 180
1668 ctatgtaggg gacctccaga cctcaggaa cctactgcaa gaggagagct accggagccg 240
1669 catcaatgag aagtctgtct ggtgctgcgg ctggcttccc tgcacaccac tgaggatcgc 300
1670 agccactgca ggccatggga actgtgtgga ctctctcata cgcaaagggg ccgaggtgga 360
1671 cctgggtgat gtcaaggggc agactgccct gtatgtggct gtagtgaacg ggcacttggg 420
1672 gagcactgag atccttttgg aagctgtgtc tgatcccaac ggcagccggc accaccgcag 480
1673 cactcctgtg taccatgcct ytcgtgtggg tagggacgac atcctgaagg ctcttatcag 540
1674 gtatggggca gatgttgatg tcaaccatca tctgaattct gacacccggc ccccttttcc 600
1675 acggcggcta acctccttgg tggctctgtc tctatacatc agtgtgcct accataacct 660
1676 tcagtgtctc aggtctgtct tgcaggctgg ggcaaatcct gacttcaatt gcaatggccc 720
1677 tgtcaacacc caggagtctc acaggggac ccctgggtgt gtcattggtg ctgtcctgcg 780
1678 ccatggctgt gaagcagcct tcgtgagttc gttggttagg tttggagcca acctgaacct 840
1679 ggtgaagtgg gaatccctgg gcccgagggc aagaggcaga agaaagatgg atcctgaggc 900
1680 cttgcaggtc tttaaagagg ccagaagtat tcccaggacc ttgctgagtt tgtgccgggt 960
1681 ggctgtgaga agagctcttg gcaaataccg actgcattct gttccctcgc tgccgctgcc 1020
1682 agaccccata aagaagtttt tgccttatga gtacgattca catgcagtgc tgactgcaat 1080
1683 gtggaagccg atcacctgca gtgaaaactg acacagactc tggcatcctg ggaaccatgg 1140
1684 cctgtgctgc cagcttgatc cttggctgtc agtgaagaaa aaacggctgt gttctcttgg 1200
1685 actgtgattc tatctcaggt gcttggggcca tcgaacgctc cttgagtcac tgtcaactga 1260
1686 gaggcacata caaacttaat tttgttctc ttcagtctct ctgttttggg ttcttcttgg 1320
1687 caatgtgtgc agcatgggct gagcctgggt attgcctag tggggaaggc ttttttctcc 1380
1688 aggctatgca tctatttatg ttctactttt gcaatttatt gttcttttaa ggcttgatat 1440
E--> 1713 caaaacagaa agaggtttgt taagaaaaga tatagggaga aagggaattcc ggttccgtgc 1500
1715 acttctagc ctgcttttct tgcctgggtt tgtctgtcta tgcctgctgg tgcacatccc 1560
1717 ttctcttgc tgccactgtt ctattttggg agttgtcttc cgtctaagat ggcttctggg 1620

```

miscount typo
 should be 1500

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```

1719 gttctatctt attgcacaga ggtcccagaa cagtgttcat agggcaccat ctgctctgcc 1680
1721 aaggggtttc tgatgtctta cctggggat cttcagacag tggttacctt taggagaccc 1740
1723 acctggaact aaccattaag tgactgcccc cattcagatc agggaccatc ttaatagtac 1800
1725 tcaactgccag tcctcacaaag agaagatgac acgggtgctc tcttcagaca ctcccatata 1860
1727 ggaagtggga aaatgtcttg gtcacctggg ttgttcccag gctacaactt cttggtgttc 1920
1729 cactaaracc agpatatcct agttttttgg gttgactgtt cctccccac tttccttgaa 1980
W--> 1731 ncccaatgcc cttttgtttn gtttgettcc ctaaaakt                      2019

```

n is at location 2000, not 2010

F.Y.I.

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

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L:161 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:161 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:1085 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1237 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1713 M:254 E: No. of Bases conflict, LENGTH:Input:1550 Counted:1500 SEQ:24
L:1731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:1775 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1916 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1920 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:2524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L:2679 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:3416 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51